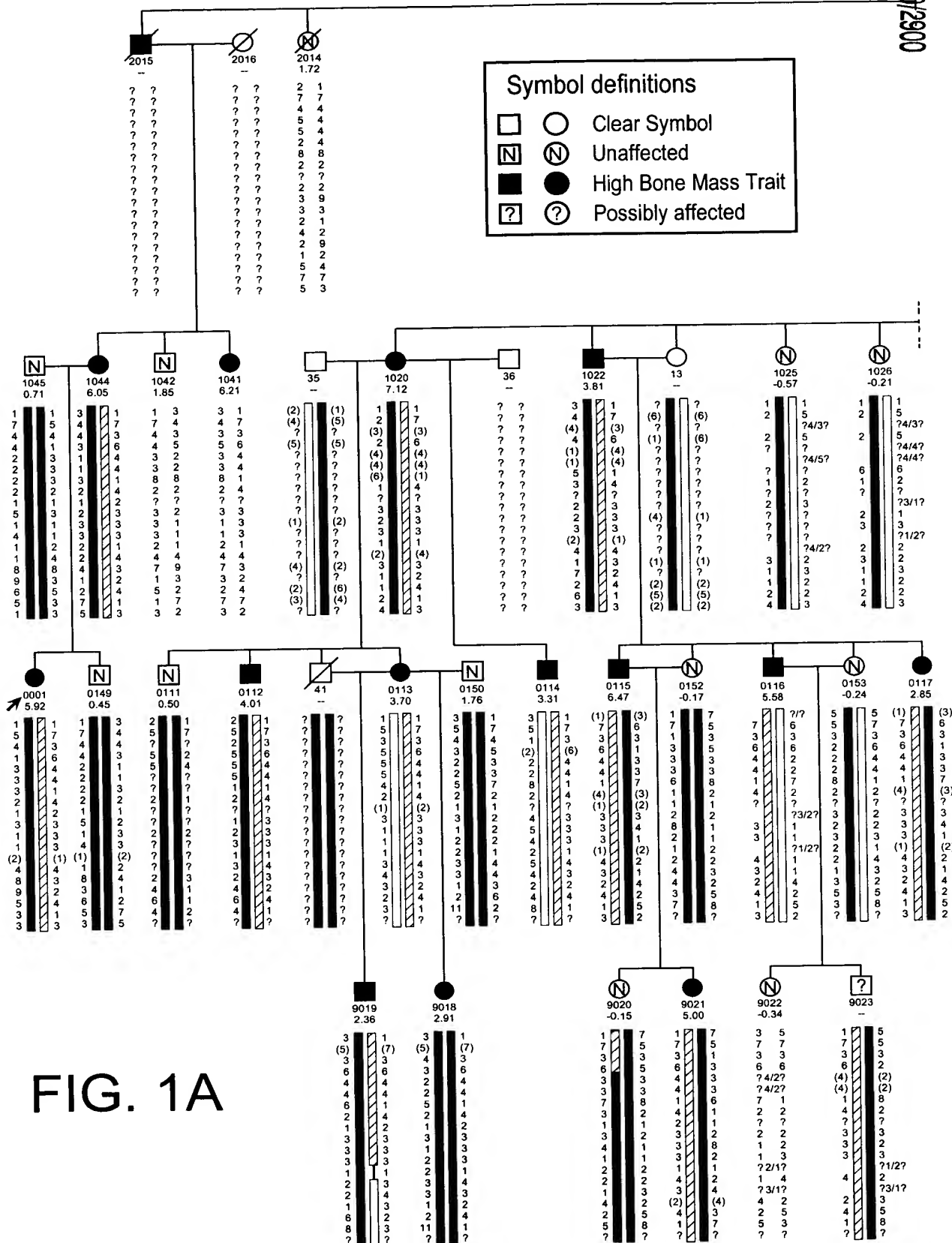




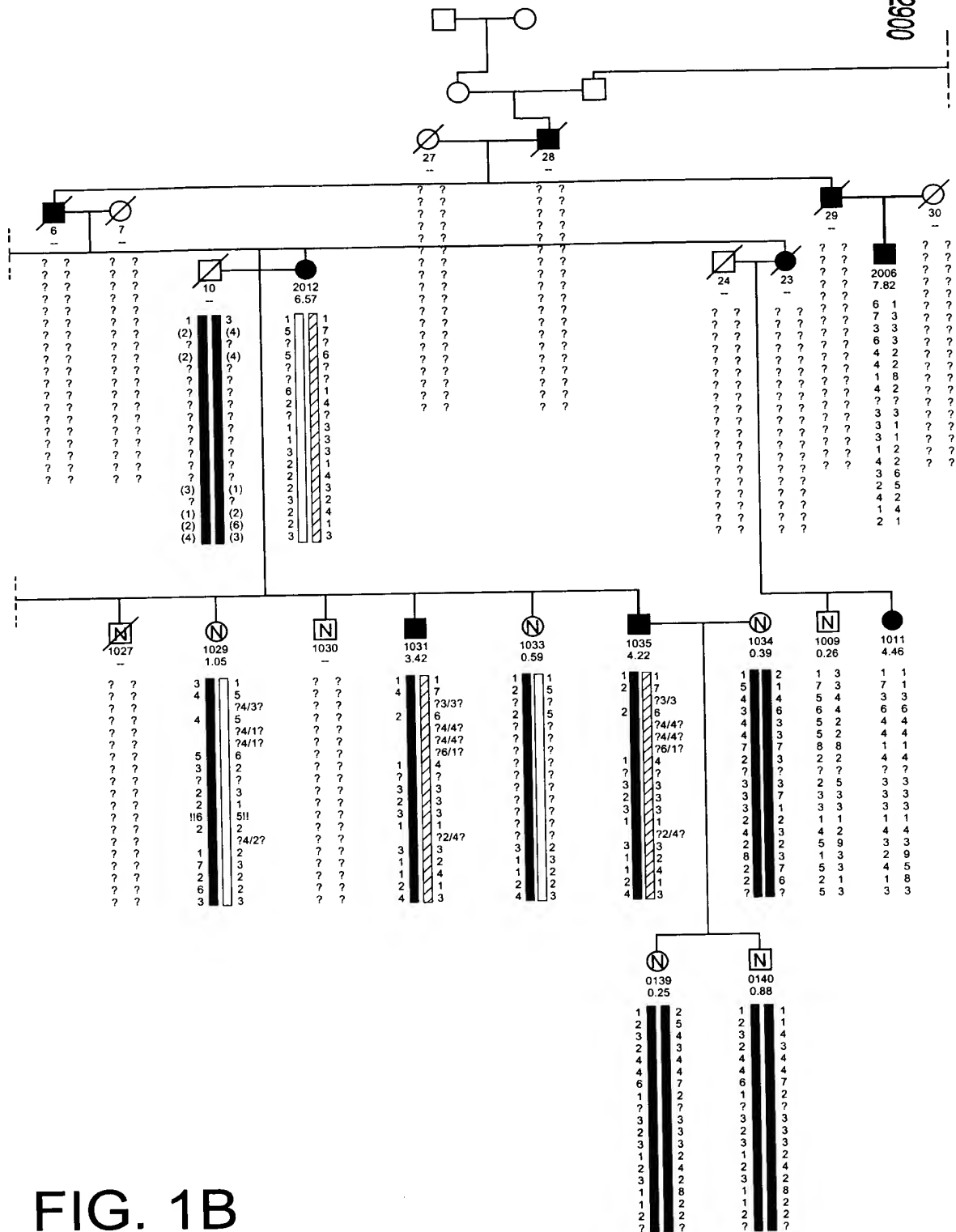
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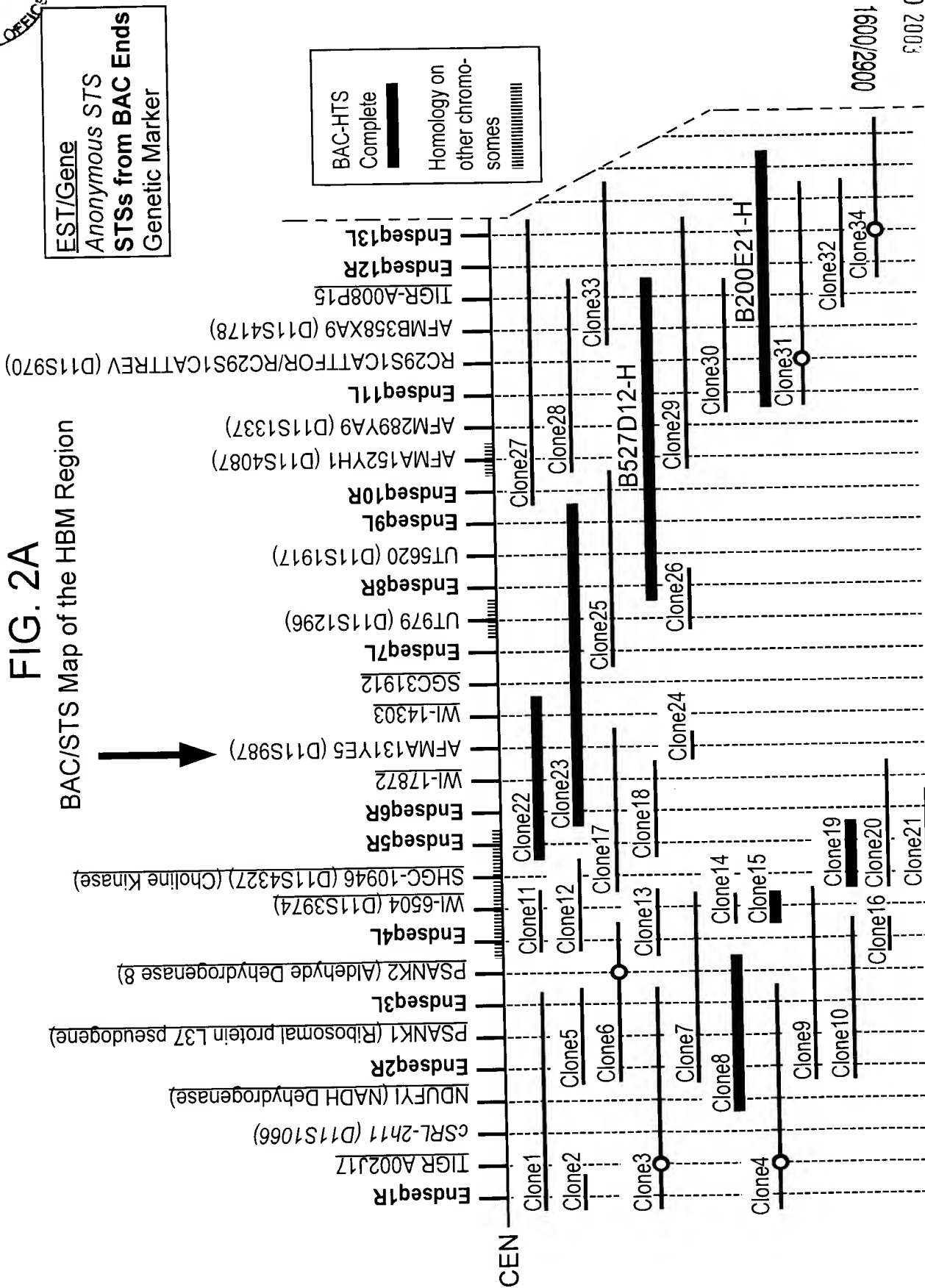
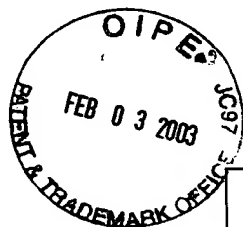
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SHEET 3 of 29

APPLN. FILING DATE: MAY 26, 2000  
 TITLE: REGULATING LIPID LEVELS VIA THE MAX1  
 OR HBM GENE  
 INVENTOR(S): JOHN P. CARULLI ET AL.  
 APPLICATION SERIAL NO: 09/578,900



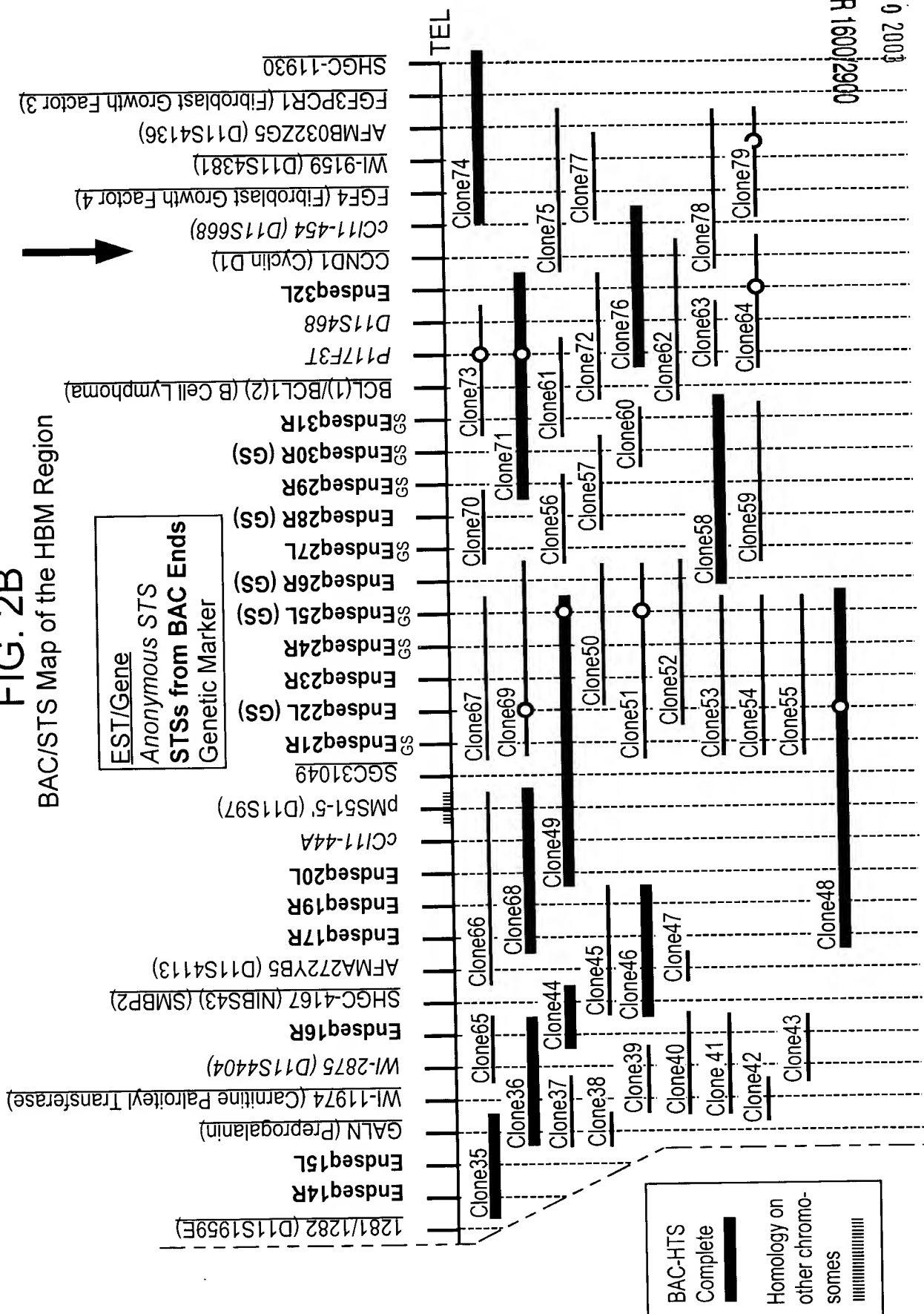
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FIG. 2B  
 BAC/STS Map of the HBM Region





**Exon 1**

ACTAAAGCGCCGCCGCCGCCATGGAGCCCGAGTGAGCGCGGGCGCG  
GGCCCGTCCGGCCGCCGACAACATGGAGGCAGCGCCGCCCGGGCCG  
CCGTGGCCGCTGCTGCTGCTGCTGCTGCTGCTGCTGGCGCTGTGCGGC  
TGCCCGGCCCGCCGCCGCC

**Exon 2 Coordinates: 527d12\_Contig308G 30944-30549**

gccccacagCCTCGCCGCTCCTGCTATTTGCCAACC GCCGGGACGTACGGC  
TGGTGGACGCCGGCGGAGTCAAGCTGGAGTCCACCATCGTGGTCAGC  
GGCCTGGAGGATGCGGCCGCGAGTGGACTTCCAGTTTCCAAGGGAGC  
CGTGTACTGGACAGACGTGAGCGAGGAGGCCATCAAGCAGACCTACCT  
GAACCAGACGGGGGCCGCCGTGCAGAACGTGGTCATCTCCGGCCTGG  
TCTCTCCCGACGGCCTCGCCTGCGACTGGGTGGGCAAGAAGCTGTACT  
GGACGGACTCAGAGACCAACCGCATCGAGGTGGCCAACCTCAATGGC  
ACATCCCGGAAGGTGCTCTTCTGGCAGGACCTTGACCAGCCGAGGGCC  
ATCGCCTTGGACCCCGCTCACGGgtaaacctgtg

... 9408 nt ...

**Exon 3 Coordinates: 527d12\_Contig308G 21141-20945**

ccccgtcacagGTACATGTACTGGACAGACTGGGGGTGAGACGCCCCGGATTG  
AGCGGGCAGGGATGGATGGCAGCACCCGGAAGATCATTGTGGACTCG  
GACATTTACTGGCCCAATGGACTGACCATCGACCTGGAGGAGCAGAAG  
CTCTACTGGGCTGACGCCAAGCTCAGCTTCATCCACCGTGCCAACCTG  
GACGGCTCGTTCCGgtaggtaccac

... 6094 nt ...

**Exon 4 Coordinates: 527d12\_Contig308G 15047-14850**

tccctgactgcagGCAGAAGGTGGTGGAGGGCAGCCTGACGCACCCCTTCGCC  
CTGACGCTCTCCGGGGACACTCTGTACTGGACAGACTGGCAGACCCGC  
TCCATCCATGCCTGCAACAAGCGCACTGGGGGGAAGAGGAAGGAGAT  
CCTGAGTGCCCTATACTACCCATGGACATCCAGGTGCTGAGCCAGGA  
CGGGCAGCCTTTCTgtgagtgcgg

... 1827 nt ...

**Exon 5 Coordinates: 527d12\_Contig308G 13220-13088**

tttctcagTCCACACTCGCTGTGAGGAGGACAATGGCGGGCTGCTCCACCTG  
TGCCTGCTGTCCCAAGCGAGCCTTTCTACACATGCGCCTGCCCCACG  
GGTGTGCAGCTGCAGGACAACGGCAGGACGTGTAAGGCAGgtgaggcggtgg  
gacg

**FIG. 3A**



... 20923 nt ...

**Exon 6 Coordinates: 527d12\_Contig309G 7705-8100**

ctccacagGAGCCGAGGAGGTGCTGCTGCTGGCCCCGGCGGACGGACCTAC  
GGAGGATCTCGCTGGACACGCCGGACTTCACCGACATCGTGCTGCAGG  
TGGACGACATCCGGCACGCCATTGCCATCGACTACGACCCGCTAGAGG  
GCTATGTCTACTGGACAGATGACGAGGTGCGGGCCATCCGCAGGGCG  
TACCTGGACGGGTCTGGGGCGCAGACGCTGGTCAACACCGAGATCAA  
CGACCCCGATGGCATCGCGGTTCGACTGGGTGGCCCCGAAACCTCTACTG  
GACCGACACGGGCACGGACCGCATCGAGGTGACGCGCCTCAACGGCA  
CCTCCCGCAAGATCCTGGTGTCTGGAGGACCTGGACGAGCCCCGAGCC  
ATCGCACTGCACCCCGTGATGGGgtaagacgggc

..... 3211 nt .....

**Exon 7 Coordinates: 527d12\_Contig309G 11311-11482**

ttcttccagCCTCATGTACTGGACAGACTGGGGAGAGAACCCTAAAATCGA  
GTGTGCCAACTTGGATGGGCAGGAGCGGCGTGTGCTGGTCAATGCCTC  
CCTCGGGTGGCCCAACGGCCTGGCCCTGGACCTGCAGGAGGGGAAGC  
TCTACTGGGGAGACGCCAAGACAGACAAGATCGAGgtgaggtcctgtgg

..... 13445 nt .....

**Exon 8 Coordinates: 527d12\_Contig309G 24927-25143**

cgtctgcagGTGATCAATGTTGATGGGACGAAGAGGCGGACCCTCCTGGA  
GGACAAGCTCCCGCACATTTTCGGGTTCACGCTGCTGGGGGACTTCAT  
CTACTGGACTGACTGGCAGCGCCGCAGCATCGAGCGGGTGCACAAGG  
TCAAGGCCAGCCGGGACGTCATATTGACCAGCTGCCCCGACCTGATGG  
GGCTCAAAGCTGTGAATGTGGCCAAGGTCGTCGgtgagtcggggggtc

....2826 nt .....

**Exon 9 Coordinates: 527d12\_Contig309G 27969-28256**

gttcgttccagGAACCAACCCGTGTGCGGACAGGAACGGGGGGTGCAGCCA  
CCTGTGCTTCTTCACACCCACGCAACCCGGTGTGGCTGCCCCATCGG  
CCTGGAGCTGCTGAGTGACATGAAGACCTGCATCGTGCCTGAGGCCTT  
CTTGGTCTTCACCAGCAGAGCCGCCATCCACAGGATCTCCCTCGAGAC  
CAATAACAACGACGTGGCCATCCCGCTCACGGGCGTCAAGGAGGCCTC  
AGCCCTGGACTTTGATGTGTCCAACAACCACATCTACTGGACAGACGT  
CAGCCTGAAGgttagcgtgggc

.....3102.....

**FIG. 3B**



**Exon 10 Coordinates: 527d12\_Contig309G 31358-31582**

cctgctgccagACCATCAGCCGCGCCTTCATGAACGGGAGCTCGGTGGAGCA  
CGTGGTGGAGTTTGGCCTTGACTACCCCGAGGGCATGGCCGTTGACTG  
GATGGGCAAGAACCTCTACTGGGCCGACACTGGGACCAACAGAATCGA  
AGTGGCGCGGCTGGACGGGCAGTTCGGCAAGTCCTCGTGTGGAGGG  
ACTTGGACAACCCGAGGTCGCTGGCCCTGGATCCCACCAAGGGgtaagtgt  
tgctgtc

.....1297 nt.....

**Exon 11 Coordinates: 527d12\_Contig309G 32879-33064**

gtgccttcagCTACATCTACTGGACCGAGTGGGGCGGCAAGCCGAGGATCG  
TGCGGGCCTTCATGGACGGGACCAACTGCATGACGCTGGTGGACAAG  
GTGGGCCCGGGCCAACGACCTCACCATTGACTACGCTGACCAGCGCCTC  
TACTGGACCGACCTGGACACCAACATGATCGAGTCGTCCAACATGCTG  
Ggtgagggccgggt

.....2069 nt.....

**Exon 12 Coordinates: 527d12\_Contig309G 35133-35454**

gtgttcagcagGTCAGGAGCGGGTTCGTGATTGCCGACGATCTCCCGCACCCG  
TTCGGTCTGACGCAGTACAGCGATTATCTACTGGACAGACTGGAAT  
CTGCACAGCATTGAGCGGGCCGACAAGACTAGCGGCCGGAACCGCAC  
CCTCATCCAGGGCCACCTGGACTTCGTGATGGACATCCTGGTGTTC  
CTCCTCCCGCCAGGATGGCCTCAATGACTGTATGCACAACAACGGGCA  
GTGTGGGCAGCTGTGCCTTGCCATCCCCGGCGGCCACCGCTGCGGCT  
GCGCCTCACACTACACCCTGGACCCCAGCAGCCGCAACTGCAGCCgtaag  
tgctcatgt

.....2006 nt.....

**Exon 13 Coordinates: 527d12\_Contig309G 37460-37659**

gcctctctaCGCCCACCACCTTCTTGCTGTTTCAGCCAGAAATCTGCCATCAG  
TCGGATGATCCCGGACGACCAGCACAGCCCGGATCTCATCCTGCCCCCT  
GCATGGACTGAGGAACGTCAAAGCCATCGACTATGACCCACTGGACAA  
GTTTCATCTACTGGGTGGATGGGCGCCAGAACATCAAGCGAGCCAAGGA  
CGACGGGACCCAGgcaggtgcctgtgg

.....6965 nt.....

**FIG. 3C**



**Exon 14 Coordinates: 527d12\_Contig309G 44624-44832**

ctttgtcttacagCCCTTTGTTTTGACCTCTCTGAGCCAAGGCCAAAACCCAGAC  
AGGCAGCCCCACGACCTCAGCATCGACATCTACAGCCGGACACTGTTC  
TGGACGTGCGAGGCCACCAATACCATCAACGTCCACAGGCTGAGCGG  
GGAAGCCATGGGGGTGGTGCTGCGTGGGGACCGCGACAAGCCCAGGG  
CCATCGTCGTCAACGCGGAGCGAGGgtaggaggccaac

.....1404 nt.....

**Exon 15 Coordinates: 527d12\_Contig309G 46236-46427**

ccaccctccgcagGTACCTGTACTTCACCAACATGCAGGACCGGGCAGCCAA  
GATCGAACGCGCAGCCCTGGACGGCACCGAGCGGAGGTCCTCTTCA  
CCACCGGCCTCATCCGCCCTGTGGCCCTGGTGGTGGACAACACACTGG  
GCAAGCTGTTCTGGGTGGACGCGGACCTGAAGCGCATTGAGAGCTGT  
GACCTGTCAGgtacgcgccccgg

.....686 nt.....

**Exon 16 Coordinates: 527d12\_Contig309G 47113-47322**

ggctgcttcagGGGCCAACCGCCTGACCCTGGAGGACGCCAACATCGTGCA  
GCCTCTGGGCCTGACCATCCTTGGCAAGCATCTCTACTGGATCGACCG  
CCAGCAGCAGATGATCGAGCGTGTGGAGAAGACCACCGGGGACAAGC  
GGACTCGCATCCAGGGCCGTGTCGCCACCTCACTGGCATCCATGCAG  
TGGAGGAAGTCAGCCTGGAGGAGTTCTgtacgtgggggc

.....3884 nt.....

**Exon 17 Coordinates: 527d12\_Contig309G 51206-51331**

ttgtctttgcagCAGCCCACCCATGTGCCCGTGACAATGGTGGCTGCTCCCACA  
TCTGTATTGCCAAGGGTGATGGGACACCACGGTGCTCATGCCCAGTCC  
ACCTCGTGCTCCTGCAGAACCTGCTGACCTGTGGAGgtaggtgtgacctaggtgc

....3905 nt.....

**Exon 18 Coordinates: 527d12\_Contig309G 55236-55472**

gttctctctgtcctccccagAGCCGCCCACCTGCTCCCCGGACCAGTTTGCATGT  
GCCACAGGGGAGATCGACTGTATCCCCGGGGCCTGGCGCTGTGACGG  
CTTTCCCGAGTGCGATGACCAGAGCGACGAGGAGGGCTGCCCCGTGT  
GCTCCGCCGCCAGTTCCCCTGCGCGCGGGGTCAGTGTGTGGACCTGC  
GCCTGCGCTGCGACGGCGAGGCAGACTGTCAGGACCGCTCAGACGAG  
GTGGACTGTGACGgtgaggccctcc

.....3052 nt.....

**FIG. 3D**





**Exon 19 Coordinates: 527d12\_Contig309G 58524-58634**

tctccttgagCCATCTGCCTGCCCAACCAGTTCCGGTGTGCGAGCGGCCAGT  
GTGTCCTCATCAAACAGCAGTGCGACTCCTTCCCCGACTGTATCGACG  
GCTCCGACGAGCTCATGTGTGgtgagccagctt

.....1448 nt.....

**Exon 20 Coordinates: 527d12\_Contig309G 60082-60319**

gtttgtctctggcagAAATCACCAAGCCGCCCTCAGACGACAGCCCCGGCCCCACA  
GCAGTGCCATCGGGCCCGTCATTGGCATCATCCTCTCTCTCTTCGTCAT  
GGGTGGTGTCTATTTTGTGTGCCAGCGCGTGGTGTGCCAGCGCTATGC  
GGGGGCCAACGGGGCCCTTCCCGCACGAGTATGTCAGCGGGACCCCGC  
ACGTGCCCCCTCAATTTTCATAGCCCCGGGCGGTTCCAGCATGGCCCCCT  
TCACAGgtaaggagcctgagatatgaa

....1095 nt.....

**Exon 21 Coordinates: 527d12\_Contig309G 61414-61552**

cttcctgcccagGCATCGCATGCGGAAAGTCCATGATGAGCTCCGTGAGCCTG  
ATGGGGGGCCGGGGCGGGGTGCCCTCTACGACCGGAACCACGTCAC  
AGGGGCCTCGTCCAGCAGCTCGTCCAGCACGAAGGCCACGCTGTACCC  
GCCGgtgaggggcccggg

.....6513 nt.....

**Exon 22 Coordinates: 527d12\_Contig309G 68065-68162**

ttgctctctcagATCCTGAACCCGCCGCCCTCCCCGGCCACGGACCCCTCCC  
TGTACAACATGGACATGTTCTACTCTTCAAACATTCCGGCCCACTGCGA  
GACCGTACAGgtaggacatcccctgag

.....2273 nt.....

**FIG. 3E**



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Exon 23 Coordinates: 527d12\_Contig309G 70435-70901

tcaaacattccggccactgcgagaccgtacagGCCCTACATCATTTCGAGGAATGGCGCCCC  
CGACGACGCCCTGCAGCACCGACGTGTGTGACAGCGACTACAGCGCC  
AGCCGCTGGAAGGCCAGCAAGTACTACCTGGATTTGAACTCGGACTCA  
GACCCCTATCCACCCCCACCCACGCCCCACAGCCAGTACCTGTCGGCG  
GAGGACAGCTGCCCCGCCCTCGCCCCGCCACCGAGAGGAGCTACTTCCAT  
CTCTTCCCCGCCCCCTCCGTCCCCCTGCACGGACTCATCCTGACCTCGGC  
CGGGCCACTCTGGCTTCTCTGTGCCCCTGTAAATAGTTTTAAATATGAACAA  
AGAAAAAATATATTTTATGATTTAAAAAATAAATATAATTGGGATTTTAA  
AAACATGAGAAATGTGAACTGTGATGGGGTGGGCAGGGCTGGGAGAACTT  
TGTACAGTGGAGAAATATTTATAAACTTAATTTTGTAACA

FIG. 3F



Model for a LDL Receptor-Related protein, Z<sup>max</sup>1



FIG. 4

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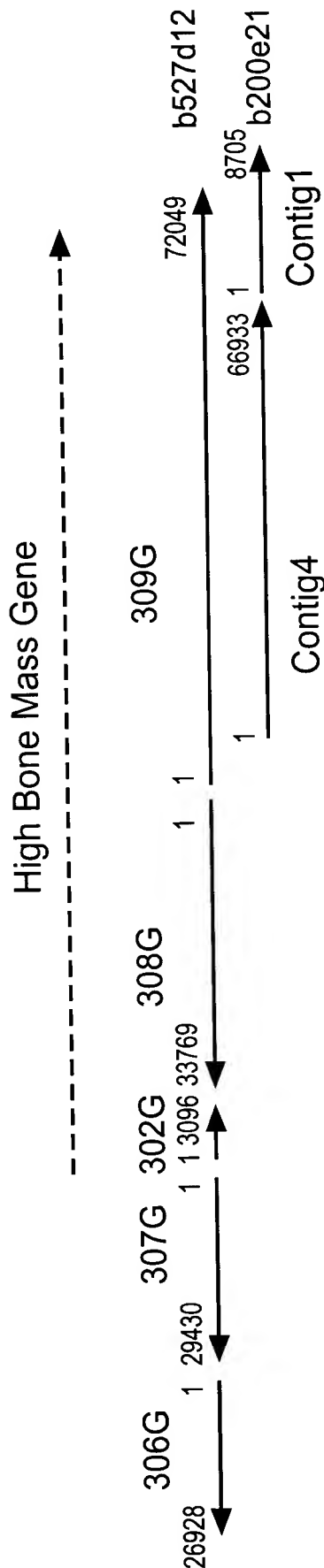


FIG. 5



FIG. 6A

1	ACTAAGCGCGCGCGCCATGGAGCCCGAGTGAGCGCGCGCGCGCGCGCGCGCC	60
61	GCCGGACAACATGGAGGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCGCT	120
1	M E A A P P G P P W P L L L L L L L	17
121	GCTGCTGCGCGCTGTGCGGCTGCCCGCGCGCGCGCGCGCGCGCGCGCGCTCCTGCTATT	180
18	L L L A L C G C P A P A A A S P L L L F	37
181	TGCCAACCGCGGACGTACGGCTGGTGACGCGCGCGCGGAGTCAAGCTGGAGTCCACCAT	240
38	A N R R D V R L V D A G G V K L E S T I	57
241	CGTGGTCAGCGGCGCTGGAGGATGCGGCGCGCAGTGAGCTTCCAGTTTCCAAAGGAGCCGT	300
58	V V S G L E D A A A V D F Q F S K G A V	77
301	GTA CTGGACAGCGTGAGCGAGGAGGCCATCAAGCAGACCTACCTGAAC CAGACGGGGC	360
78	Y W T D V S E E A I K Q T Y L N Q T G A	97
361	CGCCGTGCAGAACGTGGTCATCTCCGGCGCTGTCTCTCCGACGGCCCTCGCCTGCGACTG	420
98	A V Q N V V I S G L V S P D G L A C D W	117
421	GGTGGCAAGAGCTGTACTGGACGGA CT CAGAGACCAACCGCATCGAGGTGGCCAACCT	480
118	V G K K L Y W T D S E T N R I E V A N L	137
481	CAATGGCACATCCCGGAAGGTGCTCTTCTGGCAGGACCTTGAC CAGCGAGGGCCATCGC	540
138	N G T S R K V L F W Q D L D Q P R A I A	157
541	CTTGGACCCCGCTACGGGTACATGTACTGACAGACTGGGGTGTGAGACGCCCGGATGTA	600
158	L D P A H G Y M Y W T D W G E T P R I E	177



**FIG. 6B**

601	GCGGCAGGATGGATGGCAGCACCCGGAAGATCATTTGTGACTCGGACATTTACTGGCC	660
178	R A G M D G S T R K I I V D S D I Y W P	197
661	CAATGGACTGACCATCGACCTGGAGGAGCAGAAAGCTCTACTGGGCTGACGCCAAGCTCAG	720
198	N G L T I D L L E E Q K L Y W A D A K L S	217
721	CTTCATCCACCGTGCCAACCTGGACGGCTCGTTCCGGCAGAAAGTGGTGGAGGGCAGCCT	780
218	F I H R A N L D G S F R Q K V V E G S L	237
781	GACGCACCCCTTCGCCCTGACGCTCTCCGGGGACACTCTGTACTGGACAGACTGGCAGAC	840
238	T H P F A L T L S G D T L Y W T D W Q T	257
841	CCGCTCCATCCATGCCTGCAACAAGCGCACTGGGGGAAGAGGAAGAGATCCTGAGTGC	900
258	R S I H A C N K R T G G K R K E I L S A	277
901	CCTCTACTCACCCATGGACATCCAGGTGCTGAGCCAGGAGCGGCAGCCTTCTTCCACAC	960
278	L Y S P M D I Q V L S Q E R Q P F F H T	297
961	TCGCTGTGAGGAGGACAAATGGCGGCTGTCTCCACCTGTGCTGTCTCCCAAGCGAGCC	1020
298	R C E E D N G G C S H L C L L S P S E P	317
1021	TTTCTACACATGCGCCTGCCCCACGGGTGTGCAGCTGCAGGACAACGGCAGGACGTGTAA	1080
318	F Y T C A C P T G V Q L Q D N G R T C K	337
1081	GGCAGGAGCCGAGGAGTGCTGTCTGGCCCCGGGACGGACCTACGGAGGATCTCGCT	1140
338	A G A E E V L L L A R R T D L R R I S L	357



FIG. 6C

1141	GGACACGCCGACTTCACCGACATCGTGTGCAGGTGGACGACATCCGGCAGCCATTGC	1200
358	D T P D F T D I V L Q V D D I R H A I A	377
1201	CATCGACTACGACCCGCTAGAGGGCTATGTCTACTGGACAGATGACGAGTGGGGCCAT	1260
378	I D Y D P L E G Y V Y W T D D E V R A I	397
1261	CCGCAGGGCGTACCTGGACGGGTCTGGGGCGCAGACGCTGTCAACACCGAGATCAACGA	1320
398	R R A Y L D G S G A Q T L V N T E I N D	417
1321	CCCCGATGGCATCGCGGTCTGACTGGGTGGCCCCGAAACCTCTACTGGACCGACACGGGCAC	1380
418	P D G I A V D W V A R N L Y W T D T G T	437
1381	GGACCGCATCGAGGTGACCGGCCTCAACGGCACCTCCCGCAAGATCCTGGTGTGCGGAGGA	1440
438	D R I E V T R L N G T S R K I L V S E D	457
1441	CCTGGACGAGCCCCGAGCCATCGCACTGCACCCCGTGTGGGCTCATGTACTGGACAGA	1500
458	L D E P R A I A L H P V M G L M Y W T D	477
1501	CTGGGGAGAGAACCTAAATCGAGTGTGCCAACTTGGATGGGAGGAGCGGCGTGTGCT	1560
478	W G E N P K I E C A N L D G Q E R R V L	497
1561	GGTCAATGCCCTCCCTCGGGTGGCCCAACGGCCTGGCCCTGGACCTGCAGGAGGGGAAGCT	1620
498	V N A S L G W P N G L A L D L Q E G K L	517
1621	CTACTGGGAGACGCCAAGACAGACAAGATCGAGGTGATCAATGTTGATGGGACGAAGAG	1680
518	Y W G D A K T D K I E V I N V D G T K R	537

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FIG. 6D

1681	GCGGACCCCTCCTGGAGGACAAGCTCCCGCACATTTTCGGGTTTACGCTGCTGGGGACTT	1740
538	R T L L E D K L P H I F G F T L L G D F	557
1741	CATCTACTGGACTGACTGGCAGCGCGCCGAGCATCGAGCGGGTGACAAAGTCAAGGCCAG	1800
558	I Y W T D W Q R R S I E R V H K V K A S	577
1801	CCGGACGTCATCATTTGACCAGCTGCCCGACCTGATGGGCTCAAAGCTGTGAATGTGGC	1860
578	R D V I I D Q L P D L M G L K A V N V A	597
1861	CAAGGTCGTCGGAACCAACCCGTGTGCGGACAGGAACGGGGGTGCAGCCACCTGTGCTT	1920
598	K V V G T N P C A D R N G G C S H L C F	617
1921	CTTCACACCCACGCAACCCGGTGTGGTGCCCATCGGCCCTGGAGCTGCTGAGTGACAT	1980
618	F T P H A T R C G C P I G L E L L S D M	637
1981	GAAGACCTGCATCGTGCCCTGAGGCCTTCTTGGTCTTCAACCAGCAGAGCCGCCATCCACAG	2040
638	K T C I V P E A F L V F T S R A A I H R	657
2041	GATCTCCCTCGAGACCAATAACAACGACGTGGCCATCCCGCTCACGGGCGTCAAGGAGGC	2100
658	I S L E T N N N D V A I P L T G V K E A	677
2101	CTCAGCCCTGGACTTTGATGTGTCCAACAACACATCTACTGGACAGACGTCAGCCTGAA	2160
678	S A L D F D V S N N H I Y W T D V S L K	697
2161	GACCATCAGCCGCGCTTCATGAACGGGAGCTCGGTGGAGCACGTTGGAGTTTGGCCT	2220
698	T I S R A F M N G S S V E H V V E F G L	717

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FIG. 6E

2221	TGACTACCCCGAGGCGATGGCCGTTGACTGGGCAAGAACCTCTACTGGGCCGACAC	2280
718	D Y P E G M A V D W M G K N L Y W A D T	737
2281	TGGGACCAACAGAAATCGAAGTGGCGCGGCTGGACGGGAGTTCCGGCAAGTCTCTGTGTG	2340
738	G T N R I E V A R L D G Q F R Q V L V W	757
2341	GAGGACTTGGACAACCCGAGGTCGCTGGCCCTGGATCCACCAAGGCTACATCTACTG	2400
758	R D L D N P R S L A L D P T K G Y I Y W	777
2401	GACCGAGTGGGGCGGCAAGCCGAGGATCGTGGGCCCTTCATGGACGGGACCAACTGCAT	2460
778	T E W G G K P R I V R A F M D G T N C M	797
2461	GACGCTGGTGACAAGGTGGGGCGGCAACGACCTCACCATTTGACTACGCTGACCAGCG	2520
798	T L V D K V G R A N D L T I D Y A D Q R	817
2521	CCTCTACTGGACCGACCTGGACACCAACATGATCGAGTCGTCCAAACATGCTGGGTCAGGA	2580
818	L Y W T D L D T N M I E S S N M L G Q E	837
2581	GCGGTCGTGATTGCCGACGATCTCCCGCACCCGTTCCGGTCTGACGCGAGTACAGCGATTA	2640
838	R V V I A D D L P H P F G L T Q Y S D Y	857
2641	TATCTACTGGACAGACTGGAATCTGCACAGCATTTGACGGGGCCGACAAAGACTAGCGGCCG	2700
858	I Y W T D W N L H S I E R A D K T S G R	877
2701	GAACCGCACCCCTCATCCAGGGCCACCTGGACTTCGTGATGGACATCCTGGTGTTCACCTC	2760
878	N R T L I Q G H L D F V M D I L V F H S	897

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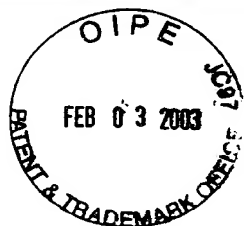
FIG. 6F

2761	CTCCCGCCAGGATGGCCTCAATGACTGTATGCACAACACGGGAGTGTTGGCAGCTGTG	2820
898	S R Q D G L N D C M H N N G Q C G Q L C	917
2821	CCTTGCCATCCCCGGGGCCACCGCTGGCGCTGCGCCTCACACTACCCCTGGACCCAG	2880
918	L A I P G G H R C G C A S H Y T L D P S	937
2881	CAGCCGCAACTGCAGCCCCGCCACCTTCTTGCTGTTCAGCCAGAAATCTGCCATCAG	2940
938	S R N C S P P T T F L L F S Q K S A I S	957
2941	TCGGATGATCCCCGGACGACGACAGCCCGGATCTCATCTCCCTGCGCTGACTGAG	3000
958	R M I P D D Q H S P D L I L P L H G L R	977
3001	GAACGTCAAAGCCATCGACTATGACCCACTGGACAAGTTCACTACTGGGTGGATGGCG	3060
978	N V K A I D Y D P L D K F I Y W V D G R	997
3061	CCAGAACATCAAGCGAGCCAAGGACGACGGGACCCAGCCCTTTGTTTGGACCTCTCTGAG	3120
998	Q N I K R A K D D G T Q P F V L T S L S	1017
3121	CCAAGGCCAAACCCAGACAGGACGCCCCACGACCTCAGCATCGACATCTACAGCCGGAC	3180
1018	Q G Q N P D R Q P H D L S I D I Y S R T	1037
3181	ACTGTTCTGGACGTGCGAGGCCAACCAATACCATCAACGTCCACAGGCTGAGCGGGGAAGC	3240
1038	L F W T C E A T N T I N V H R L S G E A	1057
3241	CATGGGGGTGCTGCTGGGACCGCGACAAGCCAGGCCATCGTCGTCACACGCGGA	3300
1058	M G V V L R G D R D K P R A I V V N A E	1077



FIG. 6G

3301	GCGAGGGTACCTGTACTTACCAACATGCAGGACCGGGCAGCCAAAGATCGAACGGCAGC	3360
1078	R G Y L Y F T N M Q D R A A K I E R A A	1097
3361	CCTGGACGGCACCGAGCGGAGGTCTCTTCAACACCGGCTCATCCGCCCTGTGGCCCT	3420
1098	L D G T E R E V L F T T G L I R P V A L	1117
3421	GGTGGTGACAACACACTGGGCAAGCTGTCTCTGGTGACGGGACCTGAAGCGCATTGA	3480
1118	V V D N T L G K L F W V D A D L K R I E	1137
3481	GAGCTGTGACCTGTCAAGGGCCAAACCGCTGACCTGGAGGACGCCAACATCGTGCAGCC	3540
1138	S C D L S G A N R L T L E D A N I V Q P	1157
3541	TCTGGGCCCTGACCATCCTTGGCAAGCATCTCTACTGGATCGACCGCCAGCAGCAGATGAT	3600
1158	L G L T I L G K H L Y W I D R Q Q Q M I	1177
3601	CGAGCGTGTGGAGAAAGACACCGGGGACAAGCGGACTCGCATCCAGGGCCGTGTGCCCCA	3660
1178	E R V E K T T G D K R T R I Q G R V A H	1197
3661	CCTCACTGGCATCCATGCAGTGAGGAAGTCAGCCTGGAGGAGTTCTCAGCCCCACCCATG	3720
1198	L T G I H A V E E V S L E E F S A H P C	1217
3721	TGCCCCGTGACAAATGGTGGCTGCTCCACATCTGTATTGCCAAGGGTGATGGGACACCCACG	3780
1218	A R D N G G C S H I C I A K G D G T P R	1237
3781	GTGCTCATGCCAGTCCACCTCGTGTCTCCTGCAGAACCTGTGACCTGTGAGAGCCGCC	3840
1238	C S C P V H L V L L Q N L L T C G E P P	1257



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FIG. 6H

3841	CACCTGCTCCCGGACCAAGTTTGCATGTGCCACAGGGAGATCGACTGTATCCCGGGGC	3900
1258	T C S P D Q F A C A T G E I D C I P G A	1277
3901	CTGGCGCTGTGACGGCTTTCCCGAGTGGGATGACAGAGCAGAGAGGGCTGCCCGCT	3960
1278	W R C D G F P E C D D Q S D E E G C P V	1297
3961	GTGCTCCGCGCCCGAGTTCCCGCTGCGCGGGGTGAGTGTGTGACCTGCGCCTGCGCTG	4020
1298	C S A A Q F P C A R G Q C V D L R L R C	1317
4021	CGACGGCGAGGCAGACTGTGACGACCGCTCAGACGAGGTGACTGTGACGCCATCTGCCT	4080
1318	D G E A D C Q D R S D E V D C D A I C L	1337
4081	GCCCAACCAGTTCCGGTGTGCGAGCGGCCAGTGTGTCTCTCATCAACAGCAGTGCGACTC	4140
1338	P N Q F R C A S G Q C V L I K Q Q C D S	1357
4141	CTTCCCGACTGTATCGACGGCTCCGACGAGCTCATGTGTGAAATCACCAGCCGCCCTC	4200
1358	F P D C I D G S D E L M C E I T K P P S	1377
4201	AGACGACAGCCCGCCACAGCAGTGCCATCGGGCCCGTCAATTGGCATCATCTCTCTCT	4260
1378	D D S P A H S S A I G P V I G I I L S L	1397
4261	CTTCGTCAATGGTGTCTATTTGTGTGCCAGCGCGTGGTGTGCCAGCGCTATGCGGG	4320
1398	F V M G G V Y F V C Q R V V C Q R Y A G	1417
4321	GGCCAACGGCCCTTCCCGCAGAGTATGTACGCGGGACCCCGCACGTGCCCTCAATT	4380
1418	A N G P F P H E Y V S G T P H V P L N F	1437



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FIG. 6I

4381	CATAGCCCCGGGGTTCCAGCATGGCCCCCTTACAGGCATCGCATCGGAAAGTCCAT	4440
1438	I A P G G S Q H G P F T G I A C G K S M	1457
4441	GATGAGCTCCGTGAGCCTGATGGGGGGCGGGGGTGGCCCTCTACGACCGGAACCA	4500
1458	M S S V S L M G G R G G V P L Y D R N H	1477
4501	CGTCACAGGGCCCTCGTCCAGCAGCTCGTCCAGCACGAAGGCCACGCTGTACCCGCCGAT	4560
1478	V T G A S S S S S T K A T L Y P P I	1497
4561	CCTGAACCCCGCCCTCCCGGCCACGGACCCCTCCCTGTACAACATGGACATGTTCTA	4620
1498	L N P P P S P A T D P S L Y N M D M F Y	1517
4621	CTCTTCAAACATTCGGGCCACTGCGAGACCGTACAGGCCCTACATCATTCGAGGAATGGC	4680
1518	S S N I P A T A R P Y R P Y I I R G M A	1537
4681	GCCCCGACGACGCCCTGCAGCACCGACGTGTGTGACAGCGACTACAGCGCCAGCCGCTG	4740
1538	P P T T P C S T D V C D S D Y S A S R W	1557
4741	GAAGGCCAGCAAGTACTACCTGGATTGAACTCGGACTCAGACCCCTATCCACCCCCACC	4800
1558	K A S K Y Y L D L N S D S D P Y P P P	1577
4801	CAGCCCCACAGCCAGTACCTGTGCGGGAGACAGCTGCCCGCCCTCGCCCCGCCACCGA	4860
1578	T P H S Q Y L S A E D S C P P S P A T E	1597
4861	GAGGAGCTACTTCCATCTCTTCCCGCCCCCTCCGTCCCCCTGCACGGACTCATCTGACC	4920
1598	R S Y F H L F P P P P S P C T D S S	1615



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FIG. 6J

4921	TCGGCCGGCCACTCTGGCTTCTCTGTGCCCCCTGTAAATAGTTTAAATATGAACAAAGA	4980
4981	AAAAAATATATTTTATGATTTAAAAATAAATAATAATTGGGATTTTAAAAACATGAGAAA	5040
5041	TGTGAACTGTGATGGGTGGCAGGGCTGGGAGAACTTTGTACAGTGGAGAAAATATTAT	5100
5101	AAACTTAATTTGTAAACA	5120

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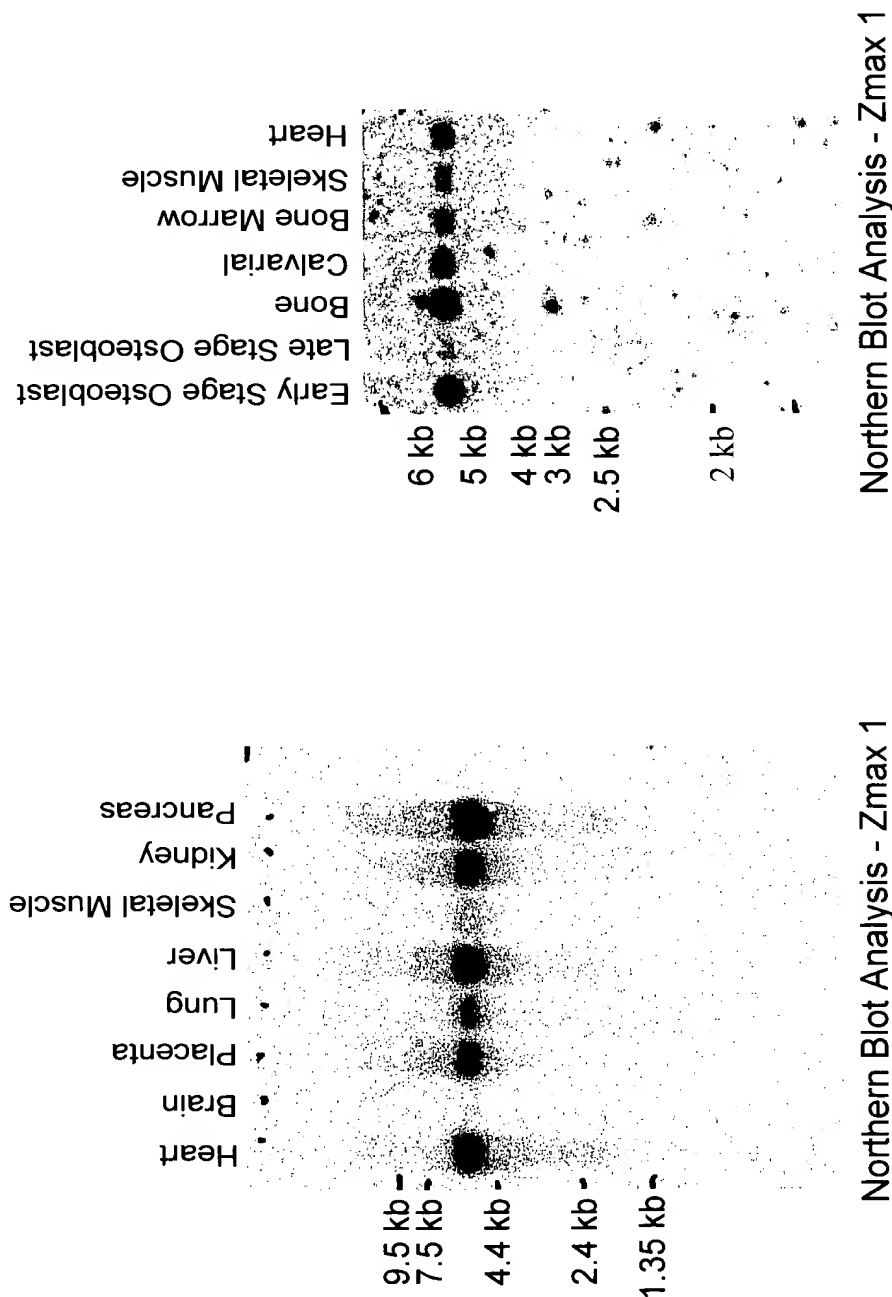


FIG. 7B

FIG. 7A

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# Zmax 1 random samples

b527d12-h\_Contig087C\_1.nt

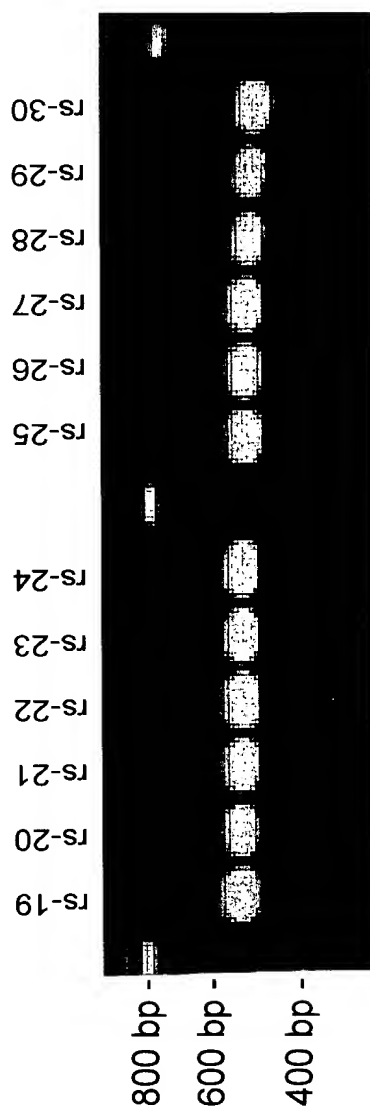


FIG. 8

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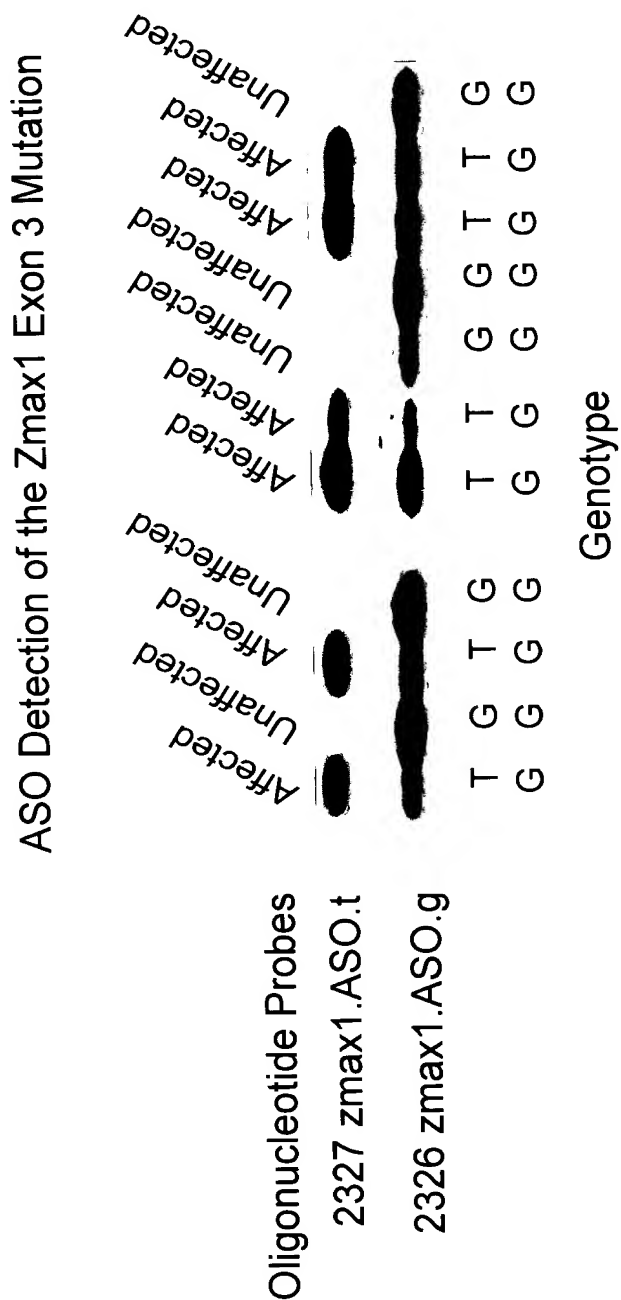


FIG. 9

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Mouse Zmax1 In situ hybridization  
100X Magnification

Antisense probe

Proliferating  
chondrocytes

Osteoblasts  
and osteoclasts

Growth  
Plate

Proximal  
aspect

Metaphysis

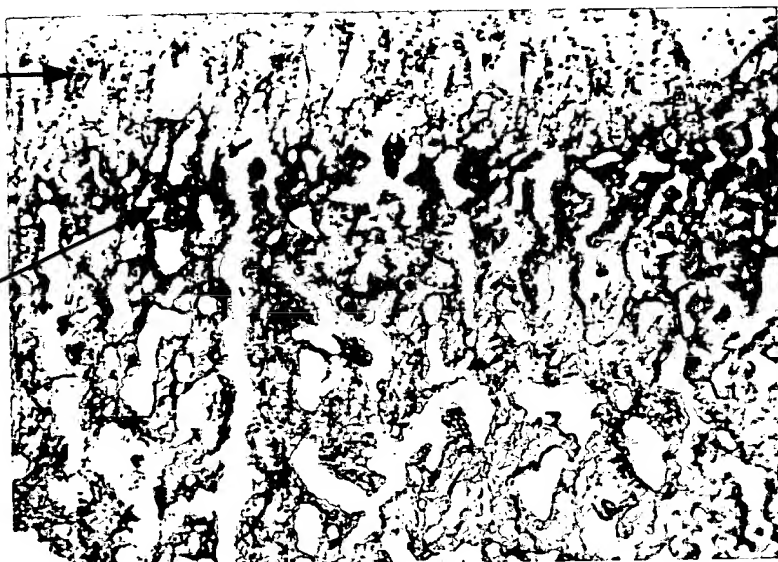


FIG. 10A

Mouse Zmax1 In situ hybridization  
100X Magnification

Sense probe



FIG. 10B

Mouse Zmax1 In situ hybridization  
400X Magnification  
Antisense probe

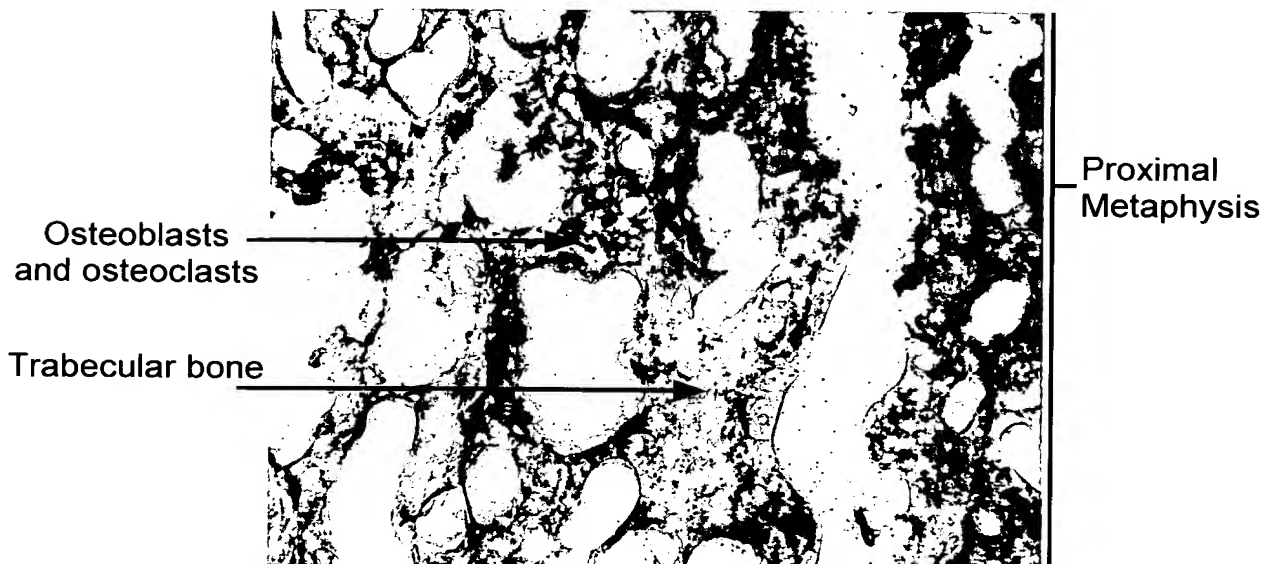


FIG. 11A

Mouse Zmax1 In situ hybridization  
400X Magnification  
Sense probe

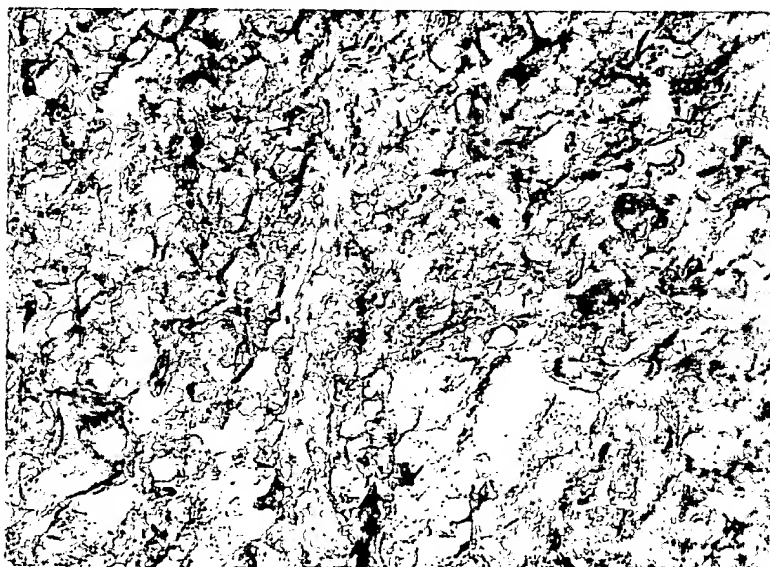


FIG. 11B

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Mouse Zmax1 In situ hybridization  
400X Magnification  
Antisense probe

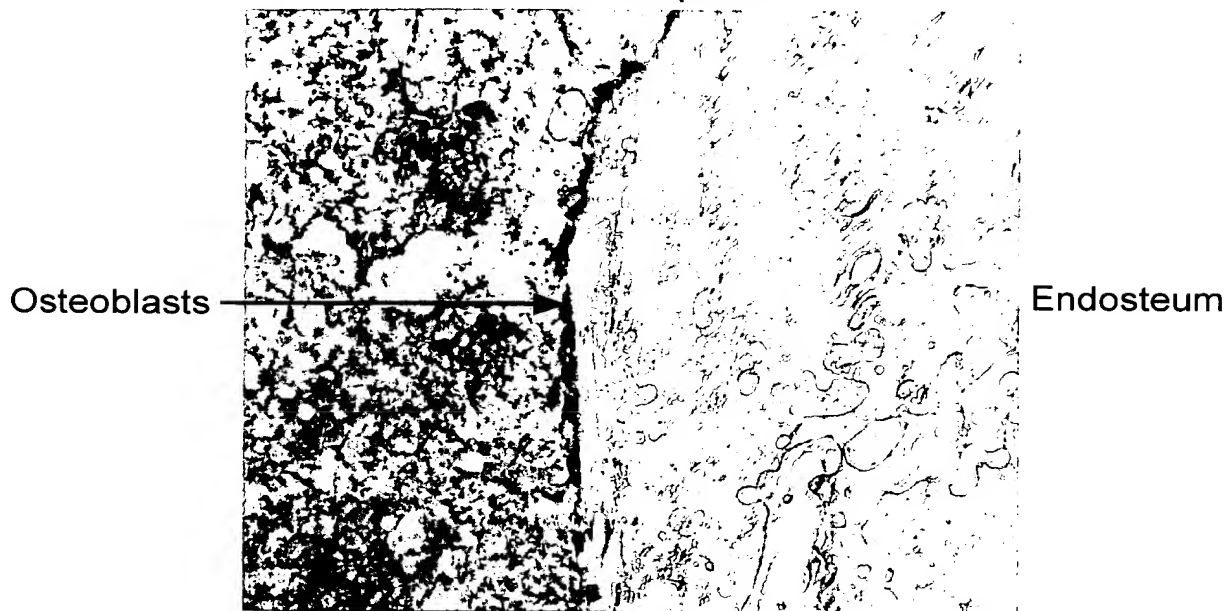


FIG. 12A

Mouse Zmax1 In situ hybridization  
400X Magnification  
Sense probe

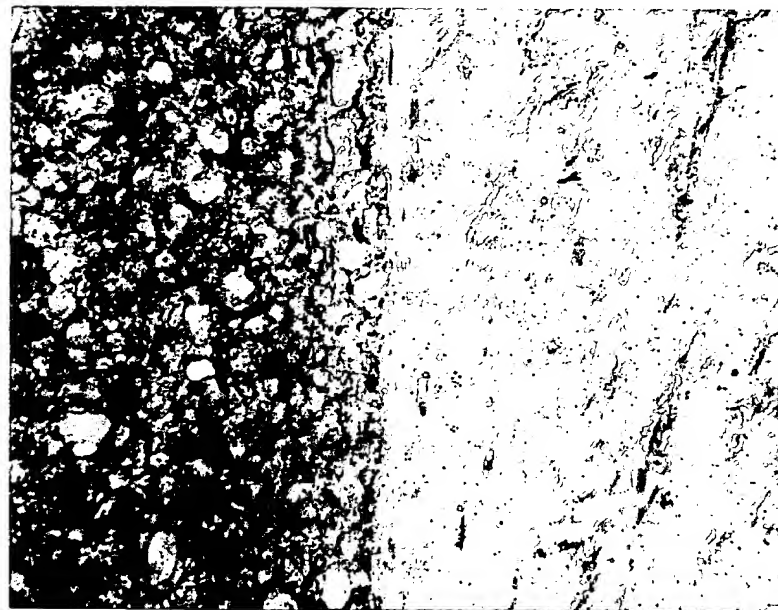


FIG. 12B

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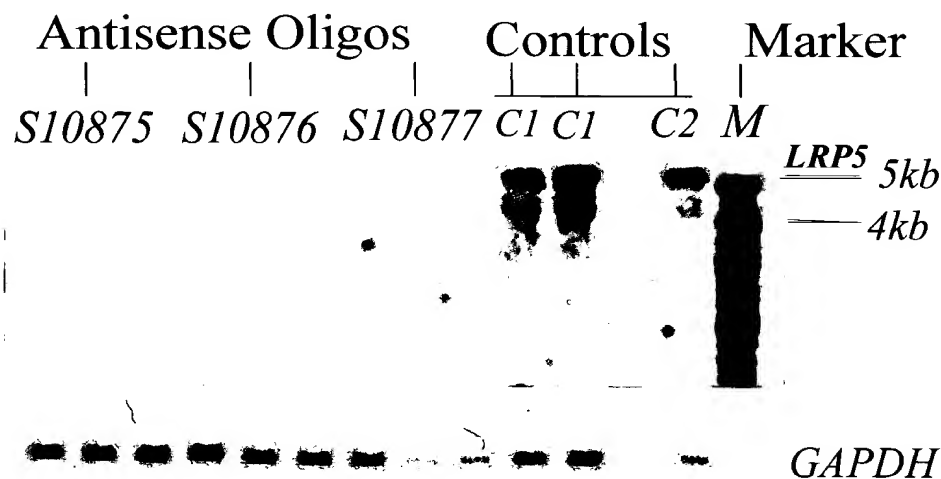
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## Antisense Inhibition of Zmax1 Expression



MC-3T3 cells

FIG. 13